

005211 070522Z60

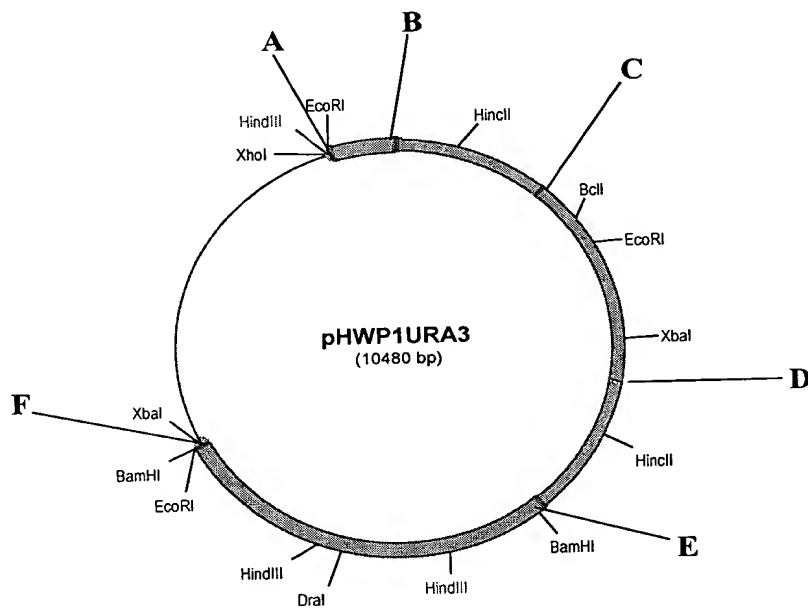


Figure 1

# Role of HWP1 in health of mice orally colonized with *C. albicans*.

Mouse type	Health	Number of mice given <i>C. albicans</i> strains of HWP1 type			
		I. <u>HWP1 HETEROZY GOTE</u>	hwp1/hwp1* homozygote	II. <u>HWP1 REVERTANT</u>	Wild type
Beige nude	ill	5	2	3	3
	not ill	2	9	1	1
	total	7	11	4	4
*P < 0.05 compared to the heterozygote, P = .058 compared to the revertant. P < .05 compared to combined heterozygote and revertant groups. Survival differences between other groups were not significant.					
Epsilon 26	ill	5	0	4	5
	Not ill	0	5	1	0
	total	5	5	5	5
*P < .01 compared to individually to the heterozygote and revertant. groups. Survival differences between other groups were not significant.					

Figure 2

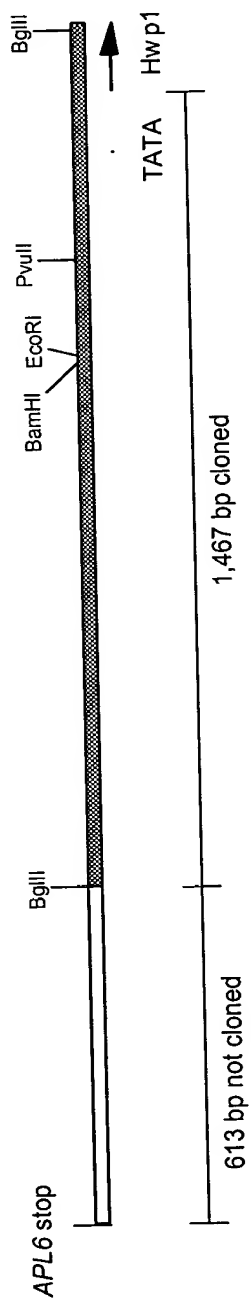


Figure 3

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171 AGAAATACAGGAAACCCCTCCAAAAAATTTTGGACCTTACACGCACATAAAATTTGCGGATAAACTTGCCATAATAAAAACTCT
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1446 ACTCGCTTTTAGTTTCGTCAATATG

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BamHI

EcoRI

PvuII

TATA box

Figure 4

0625030-1E900

TAATTCTAATAACTGATACTAAGTTTGTTCCTTTTGGGATTTCCTTTTCTTAATTTCTAATTT  
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### Figure 6

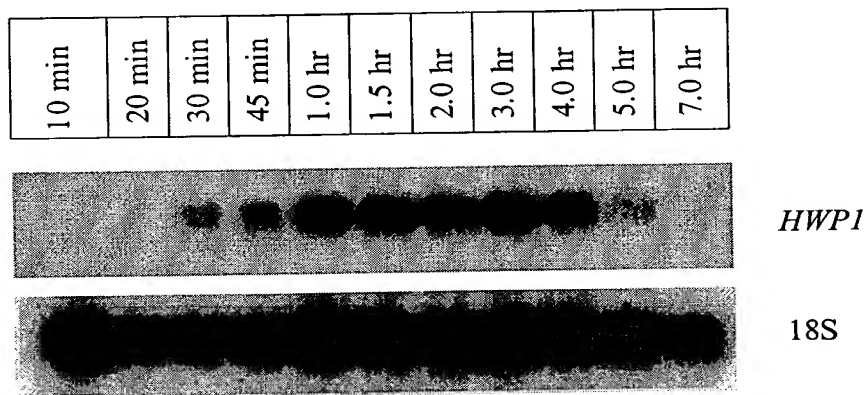
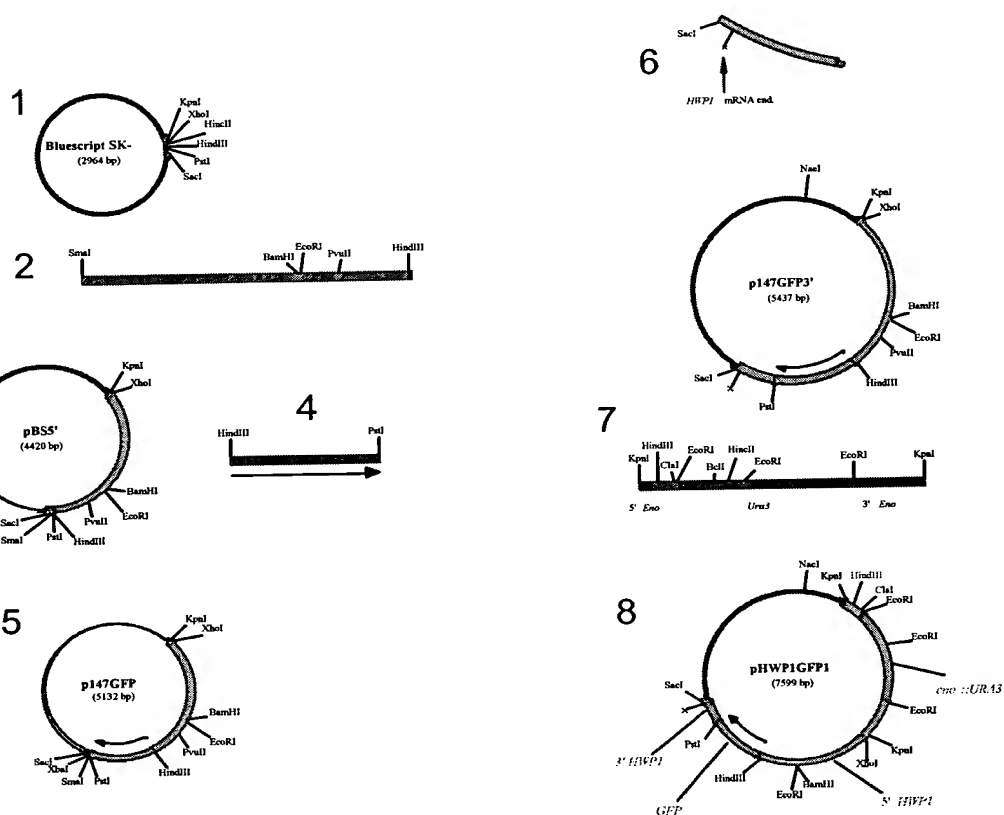


Figure 7



### Figure 8

005271 01052760

Integration of pHWP1GFP1 into the chromosome  
of *C. albicans* at the enolase locus.

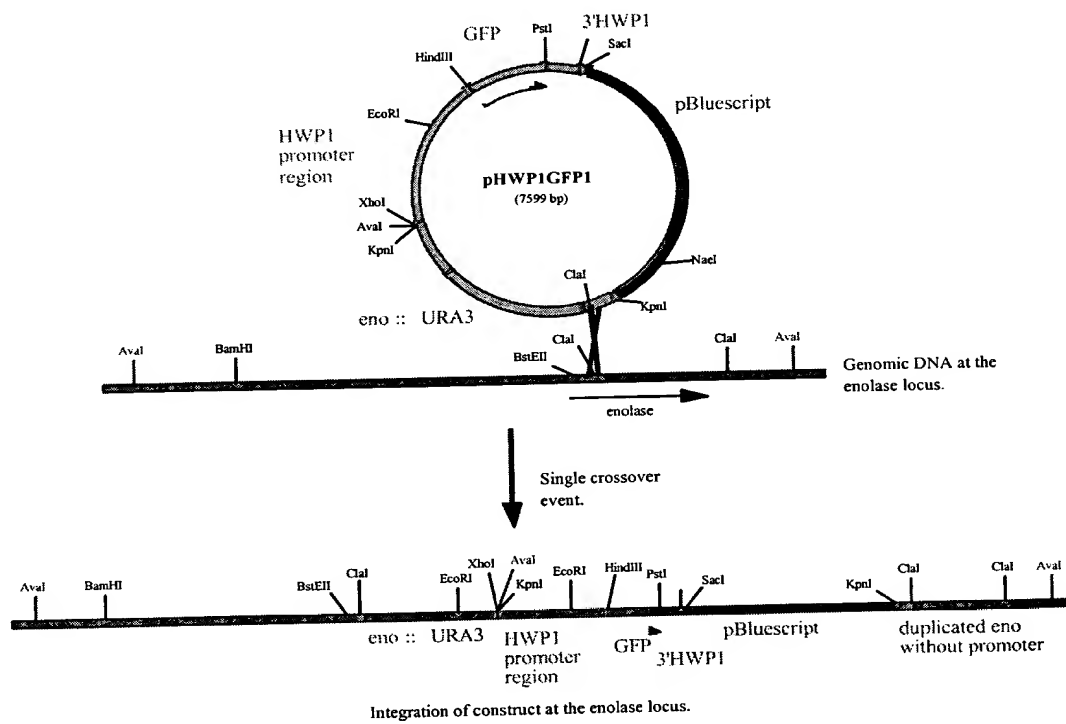


Figure 9



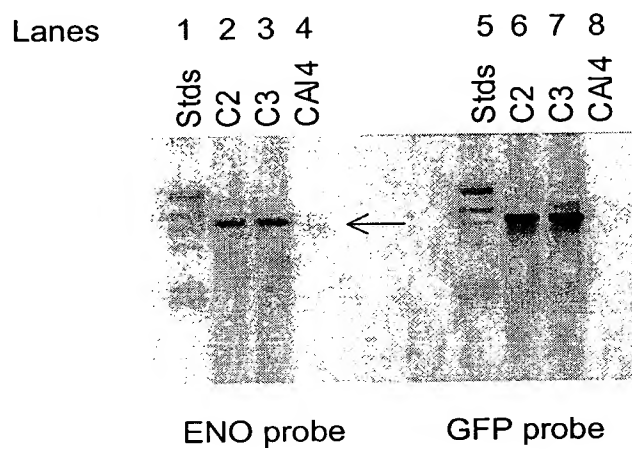


Figure 10

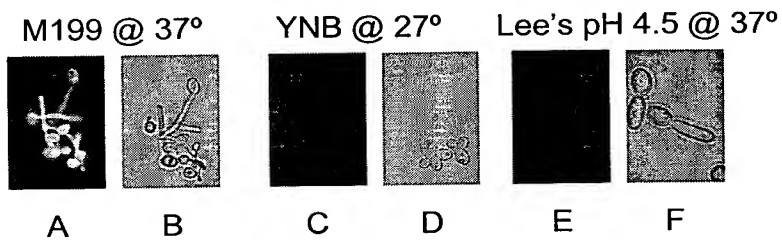


Figure 11

006277 07052260

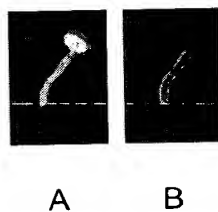


Figure 12

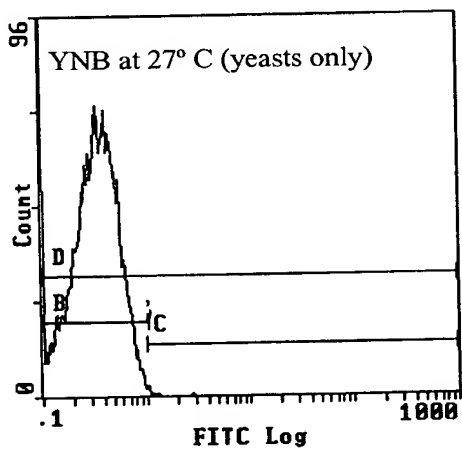


Figure 13A

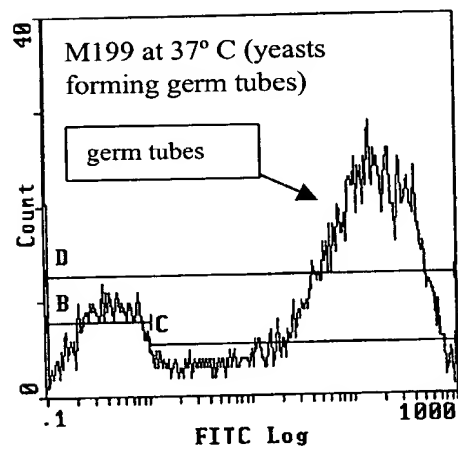
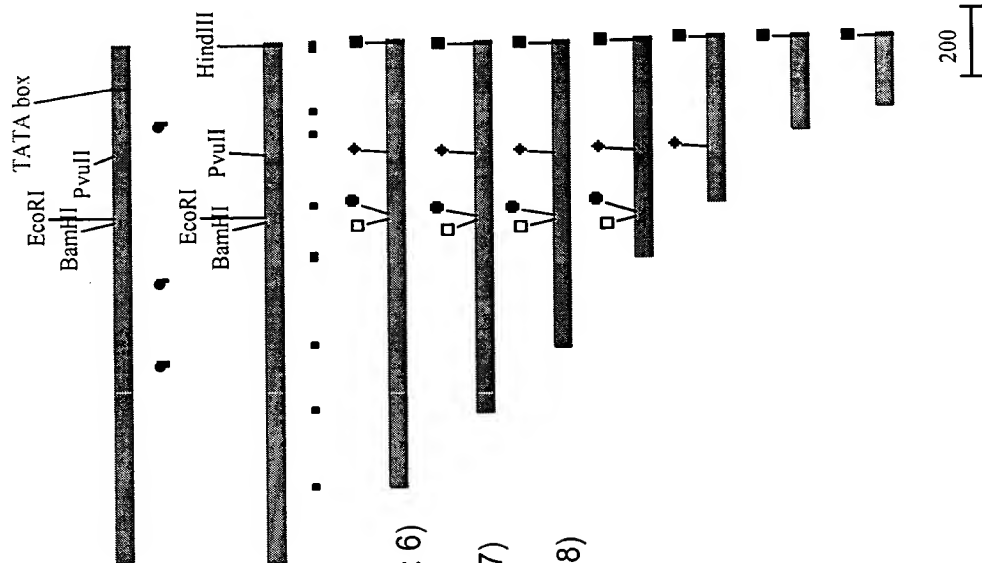


Figure 13B



5' GG CTCGAG GGA CCT TAC ACG CAC ATA AAT TGC Δ205 bp  
(SEQ ID. NO: 5)

5' GG CTCGAG CAA AAG TTA TTA GCG ATA ACC TGC Δ421 bp (SEQ ID. NO: 6)

5' GG CTCGAG GTG TAT TGT TCT CTT CAG TAC ATT Δ608 bp (SEQ ID. NO: 7)

5' GG CTCGAG CTC GAC TAA TCG ACT TTA CAT CAA Δ856 bp (SEQ ID. NO: 8)

5' GG CTCGAG ATG TCG ACT CAC AAT TCA TTG Δ1004 bp (SEQ ID. NO: 9)

5' GG CTCGAG GTT GCG ACT TTA ATA CCG TT Δ1209 bp (SEQ ID. NO: 10)

5' GG CTCGAG CAT AGC AAC TCT TGT AAA GTC Δ1271 bp (SEQ ID. NO: 11)

Figure 14A

Figure 14B

# Identification of virulence and morphogenesis factors in *C. albicans*

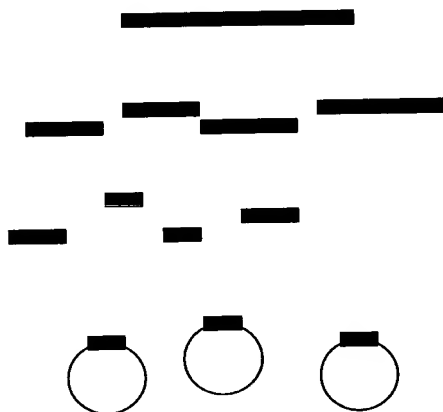
## I. STEP 1: CREATE A GENOMIC LIBRARY FOR MICROARRAY CONSTRUCTION

Prepare *C. albicans* genomic DNA.

*Sau* 3A partial digest.

Size selection of 0.5 to 2.0 kb fragments.

Clone genomic fragments into plasmid vector (pBluescript).



### A. STEP 2: Create Microarray

Transfer transformants to 96-well plates.

Perform colony PCR using universal primers.

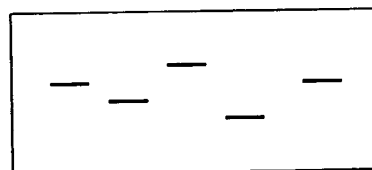
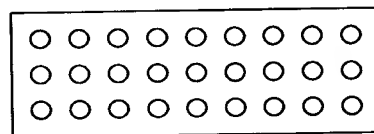
Check PCR rxns on gels and rearray positives on 96-well

Spot productive rxns on membranes.

Prepare and label cDNA from mRNA of strains with and without DNABPG

Hybridize labeled cDNA to duplicate membranes.

Go back to 96-well plates and sequence the clones of interest.



Labeled cDNA from strain; with DNABPG      dnabpg null mutant

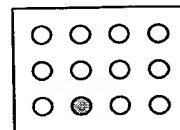
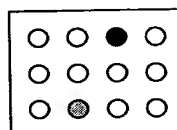
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Labeled cDNA.

wild type

mutant



→ In vivo analysis of genes.

Figure 15